line 5, "the", second occurrence, has been deleted; and line 9, "specificty" has been changed to --specificity--. Page 107, line 28, "intergrated" has been changed to --integrated--. Page 108, line 9, "aliquote" has been changed to --aliquot--; and line 20, "Contructs" has been changed to -- Constructs--. 5 Page 110, line 20, "analysis" has been changed to --analysis--; line 33, "senseitivy" has been changed to --sensitivity--; and line 35, "andGFP" has been changed to --and GFP--. Page 111, line 24, "cultur," has been changed to --culture,--; and line 26, "ligandwas" has been changed to --ligand was--. 10 Page 112, line 2 "senseitivy" has been changed to --sensitivity--. Page 133, line 6 (claim 22), "sySTEm" has been changed to --system--; and line 23 (claim 27), "Gbg" has been changed to --Gβγ --; and "STEps" has been changed to --steps--. Page 144, line 20 (claim 55), "23" has been changed to --22--. 15 Page 145, line 25 (claim 61), "23" has been changed to --22--. Page 148, line 1, "23" has been changed to --22--.

The aforementioned corrections to pages 144, 145 and 148 are evident at least, for example, from the specification at page 107, lines 1-31 (Example 12).

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Formal Drawings

Formal drawings in accordance with the guidelines of Rules 11 and 26.3(a) are submitted herewith.

L. D.

Sequence alignments of M-terminal regions of $G\alpha$ subunits and M-terminal sequences of GPA41- Gahybrid proteins.

A. Alignment of GPAl with Ga Subunits

GPA1

MGC. TVSTQTIGDESDPFLQNKRANDVIEQSLQLERQRDKNEIKLLLGAGESGKSTVLKQLKLLHQ GαS MGCLGTSKTEDQRNEEKAQREANKKIEKQLQKDKQVYRATHRLLLLGAGESGKSTIVKQMRILHV Gαi2 MGC. TVSAEDKAAERSKWIDRNLREDGEKAAKEVKLLLLGAGESGKSTIVKQWKIIHE Gαi3 MGC. TVSAEDKAAVERSKWIDRNLREDGEKAAKEVKLLLLGAGESGKSTIVKQWKIIHE
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B. GPA₄₁₋ Ga Hybrids

GPA ₄₁ - GαS MGC.TVSTQTIGDESDPFLQNKRANDVIEQSLQLLKORDKNERKLLLLGAGESGKSTIVKQMRILHV GPA ₄₁ - Gαi2	MGC.TVSTQTIGDESDPFLQNKRANDVIEQSLQLERQRDRNEVKLLLLGAGESGRSTIVRQMRIIHE GPA41 - Gai3	MGC. TVSTQTIGDESDPFLQNKRANDVIEQSLQLERQRDKNEVKLLLLGAGESGKSTIVKQMKIIHE GPA $_{41}$ - G $_{\alpha}$ 16	MGC. TVSTQTIGDESDPFLQNKRANDVIEQSLQLEKORDKNELKLLLLGPGESGKSTFIKQWRIIHG
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2/2 Coupling of Gas mutants to yeast GBy ່ວຸ. ວຸ. Š., 7. 2. 1. 1. E10R E10K ુજુડ E10A 8 _ౢౢౢౙౣౢ E10S 888 0° D1 Wt (3) (::: 0.5 2.5 0 0.2

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